

AMENDMENT

In the Specification:

Please replace paragraph [58] with the following rewritten paragraph [58]:

Q. --*S. erythraea* K97-71 contains a chromosomal deletion of the three *eryA* genes and insertion of the *xylE* gene from *Pseudomonas aeruginosa* in their place in the chromosome. To make this strain, plasmid pKOS97-49b was first constructed as follows. Two fragments flanking the *eryA* genes were PCR amplified from *S. erythraea* genomic DNA using the following primers (*Sph*I, *Hind*III, *Bam*H I, and *Eco*RI restriction sites are underlined):

eryAI left flank, forward:

5'-TTTGCATGCGGCCACGCGCACGTCGTGACC (SEQ ID NO:1),

eryAI left flank, reverse:

5'-TTAAGCTTCATATGTCCCCCTACTCGACGACCAC (SEQ ID NO:2);

eryAIII right flank, forward:

5'-TTTGGATCCGGCGGAGGGAATACATGACCACGAC (SEQ ID NO:3),

eryAIII right flank, reverse:

5'-TTTGAATTCCCGCTCGCTGAAGTCCAGGCT (SEQ ID NO:4).--

Please replace paragraph [65] with the following rewritten paragraph [65]:

Q2. --*S. erythraea* K24-1 contains a chromosomal deletion of the three *eryA* genes and insertion of the *attB* locus for the *Streptomyces* phage phiC31 from *Streptomyces lividans*, followed by the *ermE** promoter in their place. To make this strain, plasmid pKOS134-04 was first constructed as follows. The phiC31 *attB* site was inserted between the *Hind* III and *Bam*H I sites of pKOS97-49a using the following two annealed oligonucleotides:

forward: